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OM protein - protein search, using SW model

Run on: April 1, 2003, 08:45:46 ; Search time 19 Seconds
(without alignments)
995.310 Million cell updates/sec

Title: US-09-768-781-3

Perfect score: 2316

Sequence: 1 MDRYYEIPEEPNVDPESSLE.....RTRVENSEPPFETEARQSYV 449

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Listing first 45 summaries

Database : Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
 2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
 3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
 4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
 5: /cgn2_6/ptodata/1/iaa/PCUTS_COMB.pep:*
 6: /cgn2_6/ptodata/1/iaa/backfile1..pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query	Match	Length	DB ID	Description
1	164.5	7.1	129	3	US-09-056-489-46	Sequence 46, App1
2	106	4.5	451	4	US-09-134-001C-4420	Sequence 440, App1
3	100	4.3	600	4	US-09-134-001C-5655	Sequence 5655, App1
4	96.5	4.2	290	4	US-09-134-001C-4893	Sequence 4893, App1
5	95.5	4.1	255	4	US-09-154-8023	Sequence 3, App1
6	95.5	4.1	255	3	US-09-373-029-3	Sequence 3, App1
7	95.5	4.1	325	3	US-09-706-281A-18	Sequence 18, App1
8	95	4.1	325	4	US-09-097-3118	Sequence 18, App1
9	95	4.1	458	4	US-08-487-596-16	Sequence 16, App1
10	94.5	4.1	683	4	US-09-134-001C-5576	Sequence 5576, App1
11	92.5	4.0	325	1	US-08-571-525-10	Sequence 10, App1
12	92.5	4.0	325	1	US-08-672-1098-10	Sequence 10, App1
13	92.5	4.0	325	2	US-08-842-045-10	Sequence 10, App1
14	92.5	4.0	325	2	US-08-842-38-10	Sequence 10, App1
15	92.5	4.0	325	3	US-08-623-335B-10	Sequence 10, App1
16	92.5	4.0	371	1	US-08-415-818-6	Sequence 6, App1
17	92.5	4.0	371	2	US-08-894-236-6	Sequence 6, App1
18	92.5	4.0	371	2	US-08-919-624-4	Sequence 4, App1
19	92.5	4.0	371	5	PCT-US96-0144-6	Sequence 6, App1
20	91.5	4.0	378	4	US-09-259-842A-66	Sequence 66, App1
21	91.5	4.0	378	4	US-09-058-337B-66	Sequence 3507, App1
22	91.5	4.0	445	4	US-09-134-001C-3507	Sequence 3586, App1
23	91.5	4.0	739	4	US-09-151-3586	Sequence 2, App1
24	89.5	3.9	382	4	US-09-222-477-2	Sequence 3227, App1
25	89.5	3.9	443	4	US-09-134-001C-3227	Sequence 4352, App1
26	89	3.8	944	4	US-09-134-001C-4352	Sequence 4, App1
27	88.5	3.8	383	1	US-08-196-989B-4	US-09-134-001C

ALIGNMENTS

RESULT 1
US-09-058-489-46
; Sequence 46, Application US/09058489
; GENERAL INFORMATION:
; APPLICANT: Whitehead Institute for Biomedical Research
; APPLICANT: Page, David
; APPLICANT: Lahn, Bruce
; TITLE OF INVENTION: Genes in the No. 6103886-Recombinant Region of the Y Chromosome
; FILE REFERENCE: WH197-00DA
; CURRENT APPLICATION NUMBER: US/09/058, 489
; CURRENT FILING DATE: 1998-04-10
; EARLIER APPLICATION NUMBER: 60/041, 877
; EARLIER FILING DATE: 1997-04-11
; NUMBER OF SEQ ID NOS: 91
; SEQ ID NO: 46
; SOFTWARE: FastSEQ for Windows Version 3.0
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Human
US-09-058-489-46

RESULT 2
US-09-134-001C-4420
; Sequence 4420, Application US/09134001C
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC-007
; CURRENT APPLICATION NUMBER: US/09/134 001C
; CURRENT FILING DATE: 1998-08-13
; PRIORITY APPLICATION NUMBER: US 60/064, 964
; PRIORITY FILING DATE: 1997-11-08
; PRIORITY APPLICATION NUMBER: US 60/055, 779

! PRIOR FILING DATE: 1997-08-14
! NUMBER OF SEQ ID NOS: 5674
! SEQ ID NO: 4420
! LENGTH: 451
! TYPE: PRT
! ORGANISM: *Staphylococcus epidermidis*
! US-99-134-001C-4420

Query Match Score 4.6%; Score 106; DB 4; Length 451;
Best Local Similarity 17.2%; Pred. No. 0.006; Indels 146; Gaps 16;
Matches 77; Conservative 80; Mismatches 144;

Qy 30 RPTPPSILPSTPLYCGBEASALMVRVKNSTY-----RMTYTF---- 71
Db 47 KWTMISIALSLLKGALGVVQIILYMLRSQYPTQYDQEWAHYVSMCKYLIVIG 106
Qy 72 SPPMF-----SIVMQLTLLIFERDLAKD-----KPLSLPMLLILIG 108
Db 107 VPPFLPKYVQHWRINEMILRIKSTPMILLYIISLUVSNDPISILKLNPPPLIL 165
Qy 109 PVIICRLEAMKYLTKLWKKCQEQQPYVS-----LTKKMLJDGEVLIIEW 153
Db 166 -----VMYVSLIKRNLINWINQFILVIAFTFLIVIAPKSYLDEEBSRSVFK 216
Qy 154 VGHSLIRTLAMHNVAKRMSQIQAFLGSVPOLTYQIVSLSAETPLGRVLMVSLVSVT 213
Db 217 DAHRPAVIL-----AMGIVYVNLTIKQDQYDYNLLI--INIGMELYIENSRRHF 266
Qy 214 YGATLICNMIALAQIYKDDYKIR-----LGPLEVLCITIWRTLEITSLRSLILV-----F 261.
Db 267 ISVFLCLMLLPLSHIKRJIKHPJIGAMIMAIATINQPYIHHPIKJLKGNSQEVFM 326
Qy 268 SATLKLKAV-----PLVNLNLILPWEWIKRSGAQMNPNIKRSVGRGLVVLIS 314
Db 327 PSDNKAIDYALTEHPFLGSGFGI----PMIK-----ASSEIOVFNVAATNLING 373
Qy 315 VTIYAGINFSWSALQRLADLIVDKGONWGANGLAHYSVRLVNVMLVFKPFGVKV 374
Db 374 M-IFTGJ-----IGLICLTYIYMLHMLVLTFTM----- 401
Qy 375 LLNYSCHSLIALQIYLISIDFMLLF 401
Db 402 -----SITLFLPLTIVMDYIILP 422

RESULT 3
US-99-134-001C-5655
! Sequence 5655, Application US/09134001C
! Patent No. 6380370
! GENERAL INFORMATION:
! APPLICANT: Lynn Doucette-Stamm et al
! TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
! FILE REFERENCE: GTC-307
! CURRENT APPLICATION NUMBER: US/09/134,001C
! PRIOR APPLICATION NUMBER: 1998-08-13
! PRIOR FILING DATE: 1997-11-08
! PRIOR APPLICATION NUMBER: US 60/064,964
! PRIOR FILING DATE: 1997-08-14
! NUMBER OF SEQ ID NOS: 5674
! SEQ ID NO: 4593
! LENGTH: 800
! TYPE: PRT
! ORGANISM: *Staphylococcus epidermidis*
! US-99-134-001C-5655

Query Match Score 4.3%; Score 100; DB 4; Length 800;
Best Local Similarity 21.7%; Pred. No. 0.063; Indels 126; Gaps 26;

Qy 36 SLPSTPLYCGBEASALMVRVKNSTYRMTTFSPP-----MFSSIMVQLTUI 86
Db 440 SIL-TFTY-----ALYMIK-----ETFWGNNYIEKFKRKOIHEPWLPSLPAVILML 485
Qy 87 FVHDLAKDQPL-----SFMHLILGPYVRC-----LEAMKYLTKLWKKEEQEPVY 134
Db 486 I-----PVIFVPVNTGPNFYL-PATRSVSTIGAEYDAFVPHISON--HGYNLPLI 533
Qy 135 SLTRKOMLIDG-BEVILW-EVGHSLIRTLAMHNVAKRMSQ-----IQAFLGSV 181
Db 534 LSIV-----VITIGLILALVYWKVTHOILIKSASITDGYKTYREPELYSARGIRALMNN 589
Qy 182 POLTYQLYVMSLISAEPVLRGVLMVSLVSVTATLICNMIALQIYKDDYKIRLGPLEV 241
Db 590 -KLNVYIMTFLF-----VAIVYVGYLTV-GEPFHVCHIS-----SFGPLEV 632
Qy 242 CITIWRTLEIT-----SRLLILVU-----FSATLKLKAV-----PFLVLFNPLJLIFEPWIKFW 289
Db 633 LSVVTLIGLISLIFRQLRUMVWNGMIGAVTLYFIAKMPADLALTQVETITLFI 692
Qy 290 RSGAQMPN-----NIEK-NFSRQGTLVVLIS-VTLYAGINFSCWSALQLRDLVD 341
Db 693 VSFSLRLPNTERVKANLKKETFKVLSVLYMALTWSLIFY/AAQADGMPSTAKFYEDAYELT 752
Qy 342 KGQNWGHMGLHYSYVRLVENV-----MVLVPEFFGKVYVLLNY 378
Db 753 GGKN-----IVNAILGDFRALDTMFEGLVLIAGLGYITLLNY 790

RESULT 4
US-09-134-001C-4893
! Sequence 4893, Application US/09134001C
! Patent No. 6380370
! GENERAL INFORMATION:
! APPLICANT: Lynn Doucette-Stamm et al
! TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
! FILE REFERENCE: GTC-107
! CURRENT APPLICATION NUMBER: US/09/134,001C
! PRIOR APPLICATION NUMBER: US 60/064,964
! PRIOR FILING DATE: 1997-11-08
! PRIOR APPLICATION NUMBER: US 60/055,779
! PRIOR FILING DATE: 1997-08-14
! NUMBER OF SEQ ID NOS: 5674
! SEQ ID NO: 4893
! LENGTH: 290
! TYPE: PRT
! ORGANISM: *Staphylococcus epidermidis*
! US-09-134-001C-4893

Query Match Score 4.2%; Score 96.5; DB 4; Length 290;
Best Local Similarity 21.6%; Pred. No. 0.033; Indels 87; Gaps 14;
Matches 56; Conservative 42; Mismatches 74;

Qy 209 LSVTGYATLICNMIALQIYKDDYKIRLGPLEVLCITIWRTLE-----ITSLRSLILV----- 259
Db 24 LVPGVSGGTTIALLGI-----YDDFISIISGL--FSRFRWPBLKFLPILVYMLIAIGLSSQ 78
Qy 260 LFSATLKLKAVP-PLVNLNLILPWEWIKRSGAQMNPNIKRSVGRGLVVL-IVST 316
Db 79 LFNVLISQHIIPTMFLFTGILGTTIYL-----LTISNFKNSIIHWFILLGIAII 131
Qy 317 ILYAGINFSWSALQRLADRLVYDQGQNWGHM-----LRYSVRL----- 357
Db 132 VVFLALNHG-----DK-----HSGETLTLFSLIKYFAGVFASSAMLL 171
Qy 358 --VENVIMVLFVKFG-----VKVLLNYCHSLIALQIYLIAVLSI----- 220
Db 172 PGISGSFMLVFGYGTLYMAISELLHFNINALPLIILFGIVTGFM----- 220

RESULT 5
 US-09-154-802-3
 / Sequence 3, Application US/09154802
 / Patent No. 5989922
 / GENERAL INFORMATION:
 / APPLICANT: Y. Tom Tang
 / Corley, Mariah R.
 / APPLICANT: Guegler, Karl J.
 / APPLICANT: Baughn, Mariah R.
 / TITLE OF INVENTION: ATP SYNTHASE SUBUNIT HOMOLOG
 / FILE REFERENCE: PP-0596 US
 / CURRENT APPLICATION NUMBER: US/09/154,802
 / CURRENT FILING DATE: 1998-09-17
 / NUMBER OF SEQ ID NOS: 3
 / SOFTWARE: PERL Program
 / SEQ ID NO 3
 / LENGTH: 255
 / TYPE: PRT
 / ORGANISM: Homo sapiens
 / FEATURE: -
 / OTHER INFORMATION: 91006572

Query Match 4.1%; Score 95.5; DB 2; Length 255;
 Best Local Similarity 23.2%; Pred. No. 0.035; Mismatches 108; Indels 39; Gaps 10;
 Matches 58; Conservative 45; Mismatches 108; Indels 39; Gaps 10;

Qy 183 QLTQYQVSLISAREPVLGRVVLVPSLVSVTYGATLNCNMIAIQKYDDYKIRLGPI--- 237
 Db 11 QFTTRVFGLSSGLINLDTMLTSFSMSI---AVVALMLGFSI-LNDNTNILPTRLWL 66

Qy 238 -LEVLCITIWTLE---ITSRLILVLFSA---TLKIKAVPFLVNLII--- 279
 Db 67 AFSSLYFIVEKNSBOMCGLEGRLLPFPMFSLFMYILMANVSLVPSYATNAQLMWTMG 126

Qy 280 ILFEPWIKWRSQAQMPNNIKNSRV---GTLVVLISVTLYAGINFSCMS-ALQRLA 335
 Db 127 LSVAIWIGCTLG---LAHGAKEPGLFLPSGTMPLVPMELLSYMARLSLGLRG 184

Qy 336 DRDLVDKGQWGHMGHYSVRLVENVIMVLVFKPFLVNLNCHSLSIALQIITAYLISI 395
 Db 185 SNTLA---GHLVVTLAGLILNFSISMFTPALGILPLSLGIVALESAFIQAM 238

Qy 396 DFLMLFFQYL 405
 Db 239 VFTMLTCSYI 248

RESULT 7
 US-08-706-281A-18
 / Sequence 18, Application US/08706281A
 / Patent No. 6100048
 / GENERAL INFORMATION:
 / APPLICANT: Cone, Roger D
 / APPLICANT: Fan, Wei
 / APPLICANT: Boston, Bruce A
 / APPLICANT: Kesterton, Robert A
 / APPLICANT: Chen, Wenbiao
 / APPLICANT: Lu, Dongsi
 / TITLE OF INVENTION: Methods and Reagents for Discovering and
 / Using Mammalian Melanocortin Receptor Agonists and Antagonists
 / TITLE OF INVENTION: Using Mammalian Melanocortin Receptor Agonists and Antagonists
 / NUMBER OF SEQUENCES: 19
 / CORRESPONDENCE ADDRESS:
 / ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
 / STREET: 300 South Wacker Drive
 / CITY: Chicago
 / STATE: IL
 / COUNTRY: USA
 / ZIP: 60606

COMPUTER READABLE FORM:
 / MEDIUM TYPE: Floppy disk
 / COMPUTER: IBM PC compatible
 / OPERATING SYSTEM: PC-DOS/MS-DOS
 / SOFTWARE: Patent In Release #1.0, Version #1.25
 / CURRENT APPLICATION DATA:
 / APPLICATION NUMBER: US/08/706,281A
 / FILING DATE: 04-SEP-1996
 / CLASSIFICATION: 435
 / ATTORNEY/AGENT INFORMATION:
 / NAME: No. 6100048nan, Kevin E
 / REGISTRATION NUMBER: 35,303
 / REFERENCE/DOCKET NUMBER: 96,886
 / TELECOMMUNICATION INFORMATION:
 / TELEPHONE: 312-913-0001
 / TELEX: 312-913-0002

INFORMATION FOR SEQ ID NO: 18:
 / SEQUENCE CHARACTERISTICS:
 / LENGTH: 325 amino acids
 / TYPE: amino acid
 / TOPOLOGY: linear
 / MOLECULE TYPE: protein

RESULT 6
 US-09-373-029-3
 / Sequence 3, Application US/09373029
 / GENERAL INFORMATION:
 / APPLICANT: Y. Tom Tang
 / APPLICANT: Guegler, Karl J.
 / APPLICANT: Baughn, Mariah R.
 / TITLE OF INVENTION: ATP SYNTHASE SUBUNIT HOMOLOG
 / FILE REFERENCE: PP-0596 US
 / CURRENT APPLICATION NUMBER: US/09/373, 029
 / CURRENT FILING DATE: 1999-08-11
 / EARLIER FILING DATE: 1998-09-17
 / NUMBER OF SEQ ID NOS: 3
 / SOFTWARE: PERL Program
 / SEQ ID NO 3
 / LENGTH: 255
 / TYPE: PRT
 / ORGANISM: Homo sapiens
 / FEATURE: -
 / OTHER INFORMATION: 91006572

US-09-373-029-3

Query Match 4.1%; Score 95.5%; DB 3; Length 325;
 Best Local Similarity 19.5%; Pred. No. 0.051;
 Matches 64; Conservative 65; Mismatches 111; Indels 89; Gaps 17;

Qy 131 EPPVSLTRKMLIDGEVVEVIEWEVGHSHIRLAMHNRAYKRMSSQIAFLGSVPO----- 184
 Db 41 EVFLTGLVSSL---ENILV---IGAIKVNKLH-----SPMFVFGSLAVADMVSM 87

Qy 185 ----TYQYV----SLISAEVPLGRVVMVPSLVSIVTGYATLICMMLAQI-KYDD--YK 232
 Db 88 SNAWBTIVYLLNNRKLHVADTFRHIDNFSMDSMICSVVAMCSLLAAVDRITIFYA 147

Qy 233 IRLGPL----EVLCITWRTBLTSRLLLVLPE 283
 Db 148 LRYTHIMTARRSGVLIACIW-TFCUSCGIVPVIYYESKVVICHISMPTMFLPMVSLY 206

Qy 284 PWIKFWRSG----ACMP--NNIEKPNFSRVTGLVVLISVTLIYAGINFSCNSALOURLADR 337
 Db 207 HMFLLARNHVKRIASPRNTNSVRQTSMSKA---ITLML-LGIFTIVCWSPPFLHLI-- 259

Qy 338 DLVDKGQMGHMGHLYSVRLENVIMVKEFVGKVLNYCHSIALQLIAYLISIDF 397
 Db 260 -----LMISCPQNYCSCPMYSFNNYLIMCNSVID----- 291

Qy 398 MLLFFQYLHPLRSLETHVVDLHCVCCH 426
 Db 292 -----PLIVALRSQEMRRT-FKEIVCCH 313

RESULT 9
 US-08-487-596-16
 ; Sequence 16, Application US/08487596
 ; Patent No. 6440681
 ; GENERAL INFORMATION:
 ; APPLICANT: Eliot, Kathryn J.
 ; APPLICANT: Ellis, Steven B.
 ; APPLICANT: Harpold, Michael M.
 ; TITLE OF INVENTION: METHODS FOR IDENTIFYING AGONISTS AND
 ; ANTAGONISTS FOR HUMAN NEURONAL
 ; TITLE OF INVENTION: NICOTINIC ACETYLCHOLINE RECEPTORS
 ; NUMBER OF SEQUENCES: 18
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: Brown, Martin, Haller & McClain
 ; STREET: 1660 Union Street
 ; CITY: San Diego
 ; STATE: CA
 ; COUNTRY: USA
 ; ZIP: 92101
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/09/097,231
 ; FILING DATE: 12-Jun-1998
 ; CLASSIFICATION: <Unknown>
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: No. 6278038, Kevin E
 ; REGISTRATION NUMBER: 35,303
 ; REFERENCE/DOCKET NUMBER: 96, 886-C
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: 312-913-0001
 ; TELEFAX: 312-913-0002
 ; INFORMATION FOR SEQ ID NO: 18:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 325 amino acids
 ; TYPE: amino acid
 ; TOPOLOGY: linear
 ; MOLECULE TYPE: protein
 ; SEQUENCE DESCRIPTION: SEQ ID NO: 18:
 ; US-09-097-231-18

Query Match 4.1%; Score 95.5%; DB 4; Length 325;

REGISTRATION NUMBER: 33,779
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: 619-238-0599
 TELEFAX: 619-238-0062
 INFORMATION FOR SEQ ID NO: 16:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 458 amino acids
 TYPE: amino acid
 TOPOLOGY: Linear
 MOLECULE TYPE: protein
 US-08-487-596-16

Query Match 4.1%; Score 95; DB 4; Length 458;
 Best Local Similarity 25.1%; Pred. No. 0.096; Gaps 8;
 Matches 44; Conservative 35; Mismatches 64; Indels 32; Gaps 8;

Qy 73 FPMFSSIMVQTLI---FV---HRDLADKPKPSLLEMHLILG--PVIICLAMIKYKLT 123
 Db 299 YLFIFIMIVTSLTIVTVEVYINVRSSSTYHMPWKRFLQKLPKLMKDHDVDRSS 358

Qy 124 WKKBEEQEPYYV---SLTRKQMLIDGEVLIEN--EVGHSIRTLAMHNRAYKRMQIQA 176
 Db 359 PERKE-SQPVKVKRKEKKQQKQSLDGEKVLVAPLEKAADSTIRVHSRYKKEHISQVQQ 417

Qy 177 PUGSVPOLTYQVLSISAEVPLGRVVMFSLVSVTGTATCMLAIOIKYDYY 231
 Db 418 DWKFAQV-----LDRFLWFLIVSAT-GSVLIFTPAKRWLHSY 457

RESULT 10
 US-09-134-001C-5576
 Sequence 5576, Application US/09134001C
 Patent No. 638070
 GENERAL INFORMATION:
 APPLICANT: Lynn Doucette-Stamm et al
 TITLE OF INVENTION: NUCLEARIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
 TITLE OF INVENTION: EPIDERMIDS FOR DIAGNOSTICS AND THERAPEUTICS
 FILE REFERENCE: GTC-007
 CURRENT APPLICATION NUMBER: US/09/134,001C
 CURRENT FILING DATE: 1998-08-13
 PRIOR APPLICATION NUMBER: US 66/064,964
 PRIOR FILING DATE: 1997-11-08
 PRIOR FILING DATE: 1997-08-14
 NUMBER OF SEQ ID NOS: 5674
 SEQ ID NO 5576
 LENGTH: 683
 TYPE: PR
 ORGANISM: *Staphylococcus epidermidis*
 US-09-134-001C-5576

Query Match 4.1%; Score 94.5; DB 4; Length 683;
 Best Local Similarity 19.9%; Pred. No. 0.2; Gaps 19;
 Matches 82; Conservative 77; Mismatches 151; Indels 103; Gaps 19;

Qy 75 MFSQIMVQTLI---FVRLDADKPKPSLLEMHLILGIVIICLAMIKYLTURK--- 127
 Db 7 IPEFLIIVLIVVSSPFTPKP-VPLARIQ--IFLG-----MILYLTPPVERNP 55

Qy 128 EQBEPVYSLSTRKQMLIDGEVLIWEGHSIRTLAMHNRAYKRMQIQAFLGSY--POLT 185
 Db 56 DSEFFVFLVIAPLFVEG-----VNTSRVHLRKYIKVMMMA-LGIVITTVG 102

Qy 186 YQYVLSISAEVPLGRVVMPSL-----VSVTGTATCMLAII 224
 Db 103 VGLFTHIWPELPGAAATAATLCPDTAVQAITKGRKVLPKKSMITLGEESLNDAG 162

Qy 225 QIKYDDYKIRLGP-----VLCITWRLLE--ITS-----RLLI--LVLFSATKVKAVPF 272
 Db 163 II--SPKIAVGVLITGTGTSIPDIOQFLIASIGGAIIGLIGMALVRFLTRMRRGLEN 219

Qy 273 LVNINFLILFEPWIKFWRSQGAQMPNNIEKNFSRGTVLVVLISVILYAGINFSWSALQ 332

RESULT 11
 US-08-671-525B-10
 Sequence 10, Application US/08671525B
 Patent No. 5703220
 GENERAL INFORMATION:
 APPLICANT: Yamada, Tadataka
 TITLE OF INVENTION: Genes Encoding Melanocortin Receptors
 NUMBER OF SEQUENCES: 23
 CORRESPONDENCE ADDRESS:
 ADDRESS: Harness, Dickey & Pierce, P.L.C.
 STREET: P.O. Box 828
 CITY: Bloomfield Hills
 STATE: MI
 COUNTRY: US
 21IP: 48303
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: Patent In Release #1.0, Version #1.25
 CURRENT APPLICATION NUMBER: US/08671-525B
 APPLICATION NUMBER: US/08/671,525B
 FILING DATE: June 27, 1996
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: Smith, Deann F.
 REGISTRATION NUMBER: 36683
 REFERENCE/DOCKET NUMBER: 2115-000853DVB
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (810) 641-1600
 TELEFAX: (810) 641-0270
 INFORMATION FOR SEQ ID NO: 10:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 325 amino acid(s)
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 US-08-671-525B-10

Query Match 4.0%; Score 92.5; DB 1; Length 325;
 Best Local Similarity 19.1%; Pred. No. 0.11; Gaps 17;
 Matches 63; Conservative 66; Mismatches 111; Indels 89; Gaps 17;

Qy 131 EPYSLTRKMLIDGEVLIWEGHSIRTLAMHNRAYKRMQIQAFLGSVPQL--- 184
 Db 41 EVELTGLVSL---ENILV---IGAIVKRNKLH-----SPNYFVGSVSLAVADMLVSM 87

Qy 185 -----TYOLYYV---SLISAEVPLGRVVMFSLVSVTGTATCMLAIIQI-KYDD--YK 232
 Db 88 SNAVBTYIILNNKHLVATDTFTRHIDVPSMCICISVASMCSLLAVDRYITFYA 147

Qy 233 IRLGPL-----EVLCITIWTLEBITSRLLVLVLFSA---TLKLKAVPELVNFJLLE 283
 Db 148 LRYHHTMTRSGVTLACIW-TFCISCGIVIYTYESKTYVTCIATCISMFMLFPMVSLYI 206

Qy 284 PWKEWRSS---AQMP---NIEKNSRFSRGTGLVVLISVTLYAGINFSCSIALQRLADR 337
 Db 207 HMPELARNHVKRKAASPRYNSVRORTSMKGAA---ITLML-LGFIVCMSPFFHLI-- 259

Qy 338 DLVDKGQNGHMGHYSYRVLVENVIMVLYFKFGVKVLLNCHSLSIALQIJIAVLSIDF 397

Db 260 -----LMISCPQNYCSCMSYFNYLIMCNSTID----- 291.

Qy 398 MLLPQQYLPLRSLETHNVVVDLHCVCC 426

Db 292 -----PLIVALRSQEMRRT--FKEIVCCH 313

RESULT 12

US-08-672-109B-10

Sequence 10, Application US/08672109B

Patent No. 5710265

GENERAL INFORMATION:

APPLICANT: Yanada, Tadataka

TITLE OF INVENTION: Genes Encoding Melanocortin Receptors

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSE: Harness, Dickey & Pierce, P.L.C.

STREET: P.O. Box 828

CITY: Bloomfield Hills

STATE: MI

ZIP: 48303

COMPUTER READABLE FORM:

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

FILED: June 27, 1996

CLASSIFICATION: 536

ATTORNEY/AGENT INFORMATION:

NAME: Smith, Deann F.

REGISTRATION NUMBER: 36683

REFERENCE/DOCKET NUMBER: 2115-000853DVC

TELECOMMUNICATION INFORMATION:

TELEPHONE: (810)641-1600

TELEFAX: (810) 641-0270

INFORMATION FOR SEQ ID NO: 10:

SEQUENCE CHARACTERISTICS:

LENGTH: 325 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-672-109-10

MOLECULE TYPE: protein

US-08-672-109-10

Query Match 4.0%; Score 92.5%; DB 1; Length 325; Best Local Similarity 19.1%; Pred. No. 0.11; Mismatches 63; Conservatve 66; Indels 89; Gaps 17; Matches 63; Conservatve 66; Mismatches 111; Indels 89; Gaps 17;

Qy 131 EPPVSLTRKMLDGEVLEWEVGHSLTLMRNRNAYKRMQSQIAFLGSVPQL----- 184

Db 41 EFPVLTGLVSL-----ENILV--IGAIVKVNKLH-----SPHYFVYGLAVADMVLVSM 87

Qy 185 -----TYQLYV----SLISAEVPLGRVVLVFSLVSVTGCATCNMLA1QI-KYDD--YK 232

Db 88 SNAWETVTIYLNKKHLVADTFVRHIDNFDMSMCISVAVASMCSSLLIAVDRYITFYA 147

Qy 233 IRLGPL-----EVLCITIWTLEITSLILVLVFS-----TLKLKAVPFVLNFLILFE 283

Db 148 LRYHIMTARRSGVLIACIW-TFCLSGVTFVIIYYESKCVWICHLSMFFMLFEMVSLYI 206

Qy 284 PWIKFWRSG---AQMP--NNIEKNSRVSGLVVLISVTLYAGINFSCHSALQRLADR 337

Db 207 HMFLIARHIVKRIASPRVNSVRGTSMSGA---ITLML-LGTFIVSPPEFLHLI-- 259

Qy 338 DLVDKGQNGHMGHLYHSVRLVNTMVLVKEFVKVLYCHSLLJALQIJAYLISIDF 397

Db 260 -----LMISCPQNYCSCMSYFNYLIMCNSTID----- 291

Qy 398 MLLPQQYLPLRSLETHNVVVDLHCVCC 426

Db 292 -----PLIVALRSQEMRRT--FKEIVCCH 313

Qy 398 MLLPQQYLPLRSLETHNVVVDLHCVCC 426

Db 260 -----LMISCPQNYCSCMSYFNYLIMCNSTID----- 291

Qy 398 MLLPQQYLPLRSLETHNVVVDLHCVCC 426

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RESULT 14

US-08-842-238-10

Sequence 10, Application US/08842238

Patent No. 586257

GENERAL INFORMATION:

APPLICANT: Yamada, Tadataka

APPLICANT: Gantz, Ira

TITLE OF INVENTION: Genes Encoding Melanocortin Receptors

NUMBER OF SEQUENCES: 23

CORRESPONDENCE ADDRESS:

ADDRESSEE: Harness, Dickey & Pierce, P.L.C.

STREET: P.O. Box 828

CITY: Bloomfield Hills

STATE: MI

COUNTRY: US

ZIP: 48303

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/842,238

PUBLISH DATE:

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Smith, Dean F.

REGISTRATION NUMBER: 36683

REFERENCE/DOCKET NUMBER: 2115-000853DVA

TELECOMMUNICATION INFORMATION:

TELEPHONE: (810) 641-1600

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INFORMATION FOR SEQ ID NO: 10:

SEQUENCE CHARACTERISTICS:

LENGTH: 325 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-629-335B-10

Query Match 4.0%; Score 92.5; DB 2; Length 325;

Best Local Similarity 19.1%; Pred. No. 0.11; Mismatches 66; Indels 17;

Matches 63; Conservative 66; Mismatches 111; Gaps 17;

Db 131 EPYVSLTRKMLIDGEVLLIEVEGHSLTRTLAMHRYAKRMSQIQAFUGSVPOI-----184

Db 41 EVFLTIGLVSLL--ENIVL--IGAVKVKNLH-----SPMFYFGSLAVADMVLVS 87

Qy 185 -----TYQLYY----SLSAEPVPLGRVYVLMFSLVSVTYGATLGNMLAIQI-KYDD--YK 232

Db 88 SNAWETVITLNNKHVIAIDTVRHIDNVPDSMICSVASMSCLLIADEVXITIYPA 147

Qy 233 IRLGPL-----EVLCITIWRTEITSRLLLVLFSA--TLKLKAVPFLVINFLLFE 283

Db 148 LRYHHIMTARRSGVIAICW-TFCISCGVFLVYIYEVXVIIICLISMFMLFFMVSIXI 206

Qy 284 PWIKFWRSG---AQMP--NNIENKNPNSRVTGLVVLISITLYAGINFSWSALQLRLLAD 337

Db 207 HMPFLJARNHVKRIAASPRYNSVORTSMKGAA---ITMLML-LGIPIVCWSPEFLHLI-- 259

Qy 338 DLVDKCGQNWHGMGLHYSVRLVENVIMVLVKEFGVYVLLNCHSLLIAQLIAYTFD 397

Db 260 -----LMISCPQNYCSCFMSYFNMYLILIMCNSVID-----291

Qy 398 MLLFFQYLHPLRSLETHNVDYDYLHCVCH 426

Db 292 -----PLIYALRSQEMRT--FKEIVCCH 313

Search completed: April 1, 2003, 08:48:06

Job time: 22 sec/s